- a VL region having the following CDRs:
 - LC-CDR1 having the amino acid sequence of SEQ ID NO:88
 - LC-CDR2 having the amino acid sequence of SEQ ID NO:93
 - LC-CDR3 having the amino acid sequence of SEQ ID NO:95; or

(g)

- a VH region having the following CDRs:
 - HC-CDR1 having the amino acid sequence of SEQ ID NO:41
 - HC-CDR2 having the amino acid sequence of SEQ ID NO:45
 - HC-CDR3 having the amino acid sequence of SEQ ID NO:48; and
- a VL region having the following CDRs:
 - LC-CDR1 having the amino acid sequence of SEQ ID NO:88
 - LC-CDR2 having the amino acid sequence of SEQ ID NO:92
 - LC-CDR3 having the amino acid sequence of SEQ ID NO:95; or

(h)

- a VH region having the following CDRs:
 - HC-CDR1 having the amino acid sequence of SEQ ID NO:41
 - HC-CDR2 having the amino acid sequence of SEQ ID NO:45
 - HC-CDR3 having the amino acid sequence of SEQ ID NO:48; and
- a VL region having the following CDRs:
 - LC-CDR1 having the amino acid sequence of SEQ ID NO:88
 - LC-CDR2 having the amino acid sequence of SEQ ID NO:92
 - LC-CDR3 having the amino acid sequence of SEQ ID NO:97; or

(i)

- a VH region having the following CDRs:
 - HC-CDR1 having the amino acid sequence of SEQ ID NO:42
 - HC-CDR2 having the amino acid sequence of SEQ ID NO:45
 - HC-CDR3 having the amino acid sequence of SEQ ID NO:48; and
- a VL region having the following CDRs:
 - LC-CDR1 having the amino acid sequence of SEQ ID NO:88
 - LC-CDR2 having the amino acid sequence of SEQ ID NO:92
 - LC-CDR3 having the amino acid sequence of SEQ ID NO:95; or

(j)

- a VH region having the following CDRs:
 - HC-CDR1 having the amino acid sequence of SEQ ID NO:41
 - HC-CDR2 having the amino acid sequence of SEQ ID NO:44
 - HC-CDR3 having the amino acid sequence of SEQ ID NO:47; and
 - a VL region having the following CDRs:
 - LC-CDR1 having the amino acid sequence of SEQ ID NO:88

- LC-CDR2 having the amino acid sequence of SEQ ID NO:92
- LC-CDR3 having the amino acid sequence of SEQ ID NO:98.
- 39. The method according to claim 36, wherein the antigen-binding molecule comprises:
 - (a) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:24; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:74; or
 - (b) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:25; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:75; or
 - (c) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:26; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:76; or
 - (d) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:27; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:77; or
 - (e) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:28; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:78; or
 - (f) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:29; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:78; or
 - (g) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:30; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:78; or
 - (h) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:31; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:79; or
 - (i) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:32; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:79; or
 - (j) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:33; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:80; or
 - (k) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid